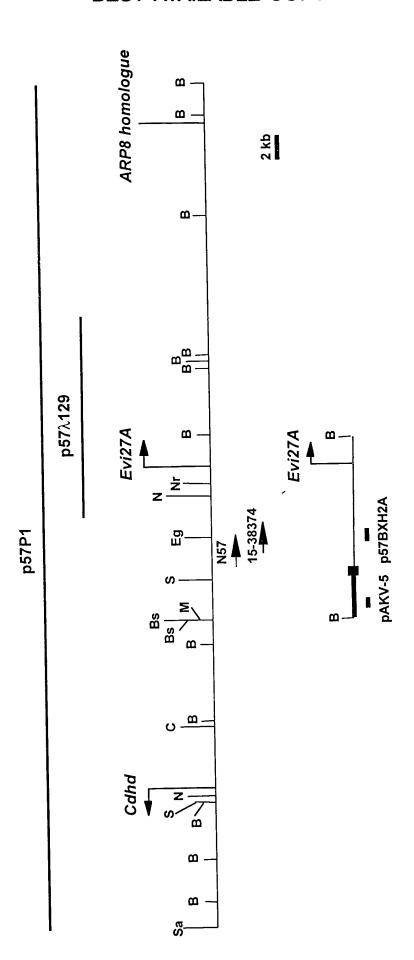
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Replacement Sheet

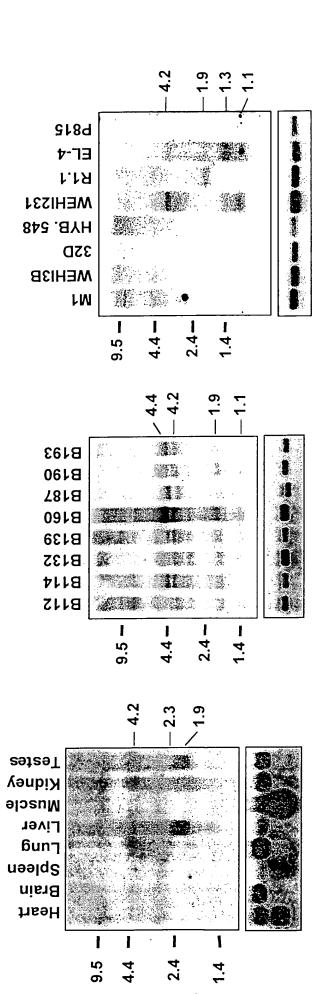


Fig. 2A

Fig. 2C

**2B** 

Fig.

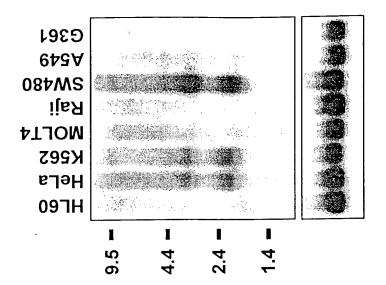


Fig. 3C

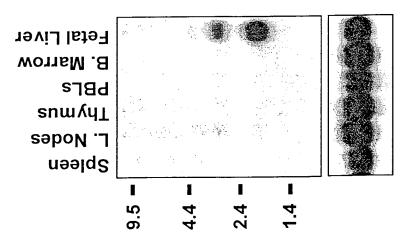


Fig. 3B

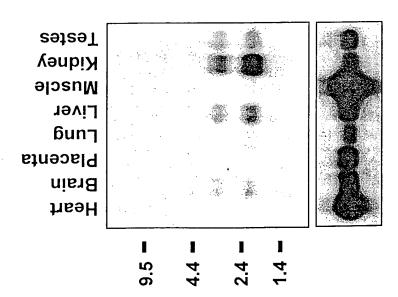
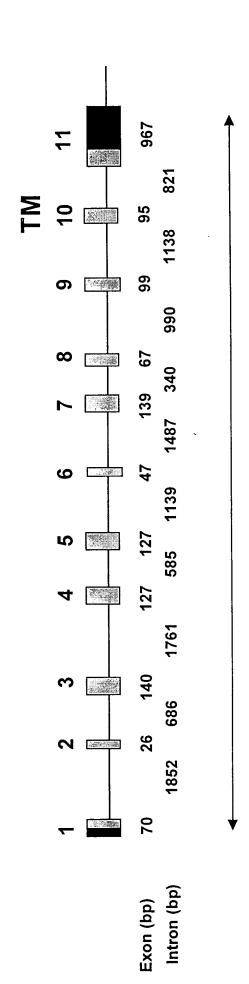


Fig. 3A

# Replacement Sheet

20	100	150 150	200	247	300	346 349	396	446 449	496	
 1 M LL V L L I L A A S C R S A L P R E P T I Q C G S E T G P S P E WM V Q H T L T P G D L R D L Q V 5 I M S L V L I S L A A L C R S A V P R E P T V Q C G S E T G P S P E WM L Q H D L I P G D L R D L R V 5 antipeptide	VAATGDYSILMNVSWVLRADASIRLLKATKICVSGKNNMNSYSCV	TEAFQSQTRPSGGKWTFSYVGFPVEL STLYLISAHNIPNANMNED STEAFQTQTRPSGGKWTFSYIGFPVEL NTVYFIGAHNIPNANMNED GNGR	TSPGCLNHVMKYKKQCTEAGSLWDPDITACKKNEKMVEVNFTT TSPGCLDHIMKYKKKCVKAGSLWDPNITACKKNEETVEVNFTT	201 NPLGNRYTILIQRDTTLGFSRVLEN - KLMRTSVAIPVTEESEGAVVQL 201 TPLGNRYMALIQHSTILGFSQVFEPHQKKQTRASVVIPVTGDSEGATVQL TM	CGSDCIRHKGTVVLCSETSAPIPPDDNRRMLGGWLPLFLV-LLVCGSDCIRHKGTVVLCPQTGVPFPLDNNKSKPGGWLPLLLSLLV	TM  297 A VWVL AAGIYL TWRQGRSTKTSFPISTMLLPLIKVLVVYPSEICFHHTVC  301 A TWVL VAGIYL MWRHERIKKTSFST-TTLLPPIKVLVVYPSEICFHHTIC	347 RFTDFLQNYCRSEVILEKWQKKKIAEMGPVQWLTTQKQAADKVVFLLPSD 350 YFTEFLQNHCRSEVILEKWQKKKIAEMGPVQWLATQKKAADKVVFLLSND GSK3	FSSQTHLHKYLVVYLG LRSQIHLHKYVVVYFR	447 G A D L K G D Y N A L S V C P Q Y H L M K D A T A F H T E L L K A T Q S M S V K K R S Q A C H D S C 450 E I D T K D D Y N A L S V C P K Y H F M K D A T A F C A E L L H V K Q Q V S A G K R S Q A C H D G C	497 S P L 499 (SEQ ID No. 7) 500 C S L 502 (SEQ ID No. 5) <b>Fig. 4</b>
Evi27 EVI27	Evi27 EVI27	Evi27 EVI27	Evi27 EVI27	Evi27 EVI27	Evi27 EVI27	Evi27 EVI27	Evi27 EVI27	Evi27 EVI27	Evi27 EVI27	Evi27 EVI27



12703 bp

Fig. 5

ILI7R EVI27	1 1	M	G	A	A	R	s	P	P	s	A	V	P	G	P	L	I M	G S	L L	L V	L L	I,	L S	L L		V A	25 10
ILI7R EVI27	26 11		A C	P -	G -	G -	A -	s -	L -	R -	L -	L -	D -	H -	R R	A S	I A		C P		Q E	P P	G	L V		- 1	50 24
ILI7R EVI27	51 25	T G	V S	K E	N T	S G	T P	C S	L P	D E	D -	s -	W W	I M	H L	P Q	R H	Ы М М	L L	T	P P	S G		P L			75 47
ILI7R EVI27	76 48				Q E			_		_			_					_			H N	Į V		W W		L L	100 72
ILI7R EVI27	101 73	Q R	T A	D D	A A	s s	I	L R	Y L	L	E K	G A	A T	E K	L	s c	y y	L T	— G	Q K	L S	N N	T F	N Q			125 97
ILI7R EVI27	126 98	L S	c c	V V	R R	F C	E N	F Y	L T	S E	K A	L F	R Q	H T	H Q	H T	R F	- P	- s	_ G	- G	R K	W W	R T	F F	T S	145 122
ILI7R EVI27	146 123	F	S I	H G	F	V P	V V	D E	P L	D N	Q T	E V	Y Y	E F	V	T G	V A	H	H N	L	P P	K N	P A	I N	P M	D N	170 147
ILI7R EVI27	171 148	G E	D	P G	N P	H S	Q M	s s	K V	N N	F F	L T	v s	P P	D G	C C	E	H D	A H	R I	M M	ĸ	Y Y	T K	T K	P K	195 172
ILI7R EVI27	196 173																									S N	220 197
ILI7R EVI27	221 198															T Q	s -	F -	P -	H -	M -	E -	N -	H H		C T	245 215
ILI7R EVI27	246 216																							L I	_		270 239
ILI7R EVI27	271 240	r T	N G	K D	L S	G -	C -	C E	R G	H A	Q T	V V	Q Q	I L	Q T	P P	F Y	F F	S P	S T	C C	L G	N S	D D	C C	L I	295 262
ILI7R EVI27	296 263	R	H H	S K	A G	T T	y y	s V	C L	P C	E P	M Q	P T	D G	T V	P P	E	P P	I L	P D	D N	Y N	M K	P S	L K	W P	320 287
ILI7R EVI27	321 288	V G	Y G	W W	F L	I P	T L	G L	I L	s L	I S	L	L L	V V	G A	s -	v -	I -	L -	L -	I -	V -	C -	M -	T T	W W	345 303
ILI7R EVI27	346 304	R	L	A V	G A	P G	G I	S Y	E L	K M	Y W	S R	D H	D E	T R	K	Y K	T K	D T	G S	L F	P S	A T	A T	D T	L L	370 328
ILI7R EVI27	371 329	I	P P	P P	]  -	L -	K -	P -	R I	K K	V V	] L	I V	I V	Y Y	S P	A	D _	H E	P I	L C	Y	V H	D H	V T	V	395 348



Fig. 6A

```
396 L K F A Q F L L T A C G T E V A L D L L E E Q A I 420
349 C Y F T E F L Q N H C R S E V I L E K W Q K K K I 373
ILI7R
EVI27
       421 | S E | A | G | V | M | T | W V | G R | Q K Q | E M V E | S | N S | K I I | V 445
ILI7R
                        QWLATQKK - - -
EVI27
       446 | L | C | S | R G T R | A | K W Q A L L | G R G A P V R L R C D 470
ILI7R
       395 | 1 | L | S | N D V N | S | V C D G T C G K S E G S -
EVI27
ILI7R
       471 H G K P V G D L F T A A M N M I L P D F K R P A C 495
EVI27
       415 P S E N S Q D L F P L A F N L F C S D L R S Q I H 439
       496 F G T Y V V C Y F S E V S C D G D
ILI7R
                                                V P D | I F G A A | 520
       440 L H K Y V V
                        V Y F R E I D T K D D Y N A L S - V C 464
EVI27
ILI7R
       521 | P R Y | P L | M D R F E E V Y F R I Q | D | L E M | F | Q P G 545
       465 P K Y H F M K - -
EVI27
       546 RMHRVGELSGDNYLRSPGGRQLRAA570
ILI7R
                                                GKRSQ-
EVI27
             LHVKQQVS
       571 L D R F R D W Q V R | Q P | D | W F E | C | E N | L | Y S A D D 595
ILI7R
                               A | C H D | G
                                                                 502
EVI27
                                                      (SEQ ID NO: 5)
ILI7R
       596 Q D A P S L D E E V F E E P L L P P G T G I V K R 620
       621 A P L V R E P G S Q A C L A I D P L V G E E G G A 645
ILI7R
ILI7R
       646 A V A K L E P H L Q P R G Q P A P Q P L H T L V L 670
ILI7R
       671 A A E E G A L V A A V E P G P L A D G A A V R L A 695
ILI7R
       696 L A G E G E A C P L L G S P G A G R N S V L F L P 720
ILI7R
       721 V D P E D S P L G S S T P M A S P D L L P E D V R 745
ILI7R
       746 E H L E G L M L S L F E Q S L S C Q A Q G G C S R 770
ILI7R
       771 PAMVLTDPHTPYEEEQRQSVQSDQG795
ILI7R
       796 Y I S R S S P Q P P E G L T E M E E E E E E Q D 820
       821 P G K P A L P L S P E D L E S L R S L Q R Q L L F 845
ILI7R
ILI7R
       846 R Q L Q K N S G W D T M G S E S E G P S A
                                                                  866
                                                      (SEQ ID NO: 9)
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Fig. 6B

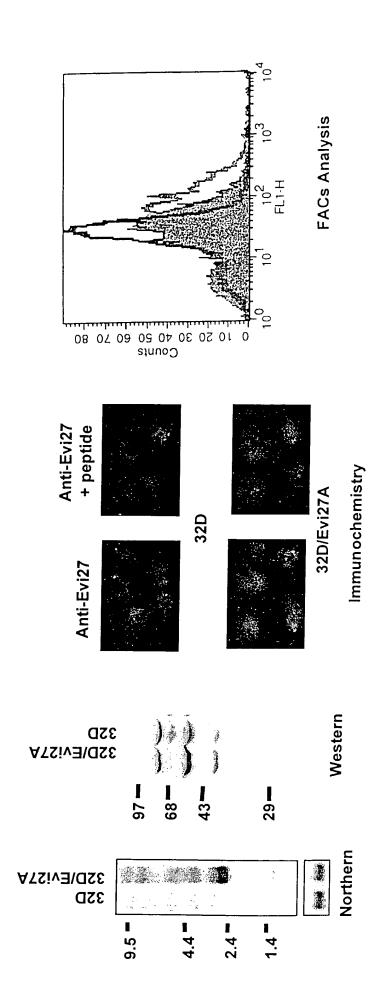


Fig. 8D

8C

Fig.

8B

Fig.

8**A** 

Fig.



=ig. 12∕

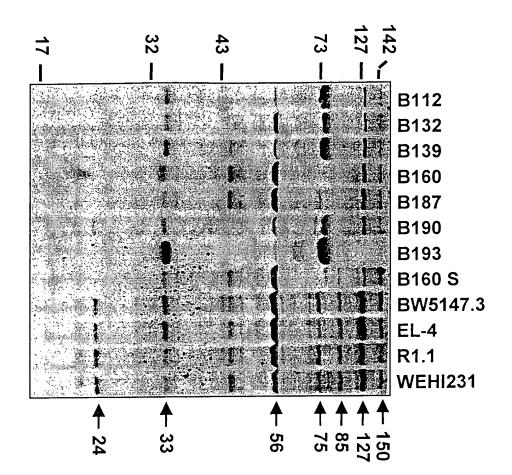
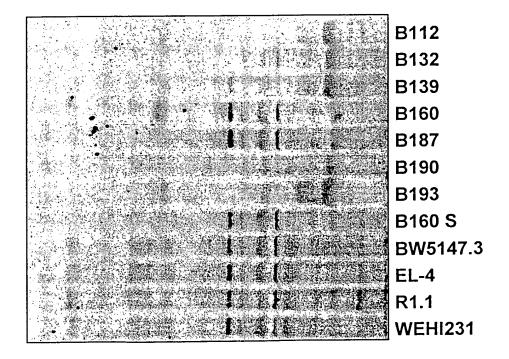


Fig. 12B



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